

# SEQUENCE LISTING



<110> Coleman et al.  
 <120> Endothelial Monocyte Activating Polypeptide III  
 <130> PF206D1  
 <140> US 08/972,301  
 <141> 1997-11-18  
 <150> US 08/483,534  
 <151> 1995-06-07  
 <160> 7  
 <170> PatentIn version 3.0  
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 <221> CDS  
 <222> (94)..(597)  
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 Glu Glu Val Ile Pro Ser Arg  
 1 5  
 ctg gat atc cgt gtg ggg aaa atc atc act gtg gag aag cac cca gat 162  
 Leu Asp Ile Arg Val Gly Lys Ile Ile Thr Val Glu Lys His Pro Asp  
 10 15 20  
 gca gac agc ctg tat gta gag aag att gac gtg ggg gaa gct gaa cca 210  
 Ala Asp Ser Leu Tyr Val Glu Lys Ile Asp Val Gly Glu Ala Glu Pro  
 25 30 35  
 cgg act gtg gtg agc ggc ctg gta cag ttc gtg ccc aag gag gaa ctg 258  
 Arg Thr Val Val Ser Gly Leu Val Gln Phe Val Pro Lys Glu Glu Leu  
 40 45 50 55  
 cag gac agg ctg gta gtg gtg ctg tgc aac ctg aaa ccc cag aag atg 306  
 Gln Asp Arg Leu Val Val Val Leu Cys Asn Leu Lys Pro Gln Lys Met  
 60 65 70  
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 Arg Gly Val Glu Ser Gln Gly Met Leu Leu Cys Ala Ser Ile Glu Gly  
 75 80 85  
 ata aac cgc cag gtt gaa cct ctg gac cct ccg gca ggc tct gct cct 402  
 Ile Asn Arg Gln Val Glu Pro Leu Asp Pro Pro Ala Gly Ser Ala Pro  
 90 95 100  
 ggt gag cac gtg ttt gtg aag ggc tat gaa aag ggc caa cca gat gag 450  
 Gly Glu His Val Phe Val Lys Gly Tyr Glu Lys Gly Gln Pro Asp Glu  
 105 110 115

gag ctc aag ccc aag aag aaa gtc ttc gag aag ttg cag gct gac ttc 498  
 Glu Leu Lys Pro Lys Lys Lys Val Phe Glu Lys Leu Gln Ala Asp Phe  
 120 125 130 135

aaa att tct gag gag tgc atc gca cag tgg aag caa acc aac ttc atg 546  
 Lys Ile Ser Glu Glu Cys Ile Ala Gln Trp Lys Gln Thr Asn Phe Met  
 140 145 150

acc aag ctg ggc tcc att tcc tgt aaa tcg ctg aaa ggg ggg aac att 594  
 Thr Lys Leu Gly Ser Ile Ser Cys Lys Ser Leu Lys Gly Gly Asn Ile  
 155 160 165

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 Ser

<210> 2  
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 35 40 45

Phe Val Pro Lys Glu Glu Leu Gln Asp Arg Leu Val Val Val Leu Cys  
 50 55 60

Asn Leu Lys Pro Gln Lys Met Arg Gly Val Glu Ser Gln Gly Met Leu  
 65 70 75 80

Leu Cys Ala Ser Ile Glu Gly Ile Asn Arg Gln Val Glu Pro Leu Asp  
 85 90 95

Pro Pro Ala Gly Ser Ala Pro Gly Glu His Val Phe Val Lys Gly Tyr  
 100 105 110

Glu Lys Gly Gln Pro Asp Glu Glu Leu Lys Pro Lys Lys Lys Val Phe  
 115 120 125

Glu Lys Leu Gln Ala Asp Phe Lys Ile Ser Glu Glu Cys Ile Ala Gln  
 130 135 140

*Sub  
G1  
cont*  
  
*J2*  
  
*cont*

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 <212> DNA  
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 <223> Contains complementary sequences to HindIII.

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 <223> Contains a BamHI restriction enzyme site.

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35 40 45  
Glu Val Asp Val Gly Glu Ile Ala Pro Arg Thr Val Val Ser Gly Leu  
50 55 60  
Val Asn His Val Pro Leu Glu Gln Met Gln Asn Arg Met Val Ile Leu  
65 70 75 80  
Leu Cys Asn Leu Lys Pro Ala Lys Met Arg Gly Val Leu Ser Gln Ala  
85 90 95  
Met Val Met Cys Ala Ser Ser Pro Glu Lys Ile Glu Ile Leu Ala Pro  
100 105 110  
Pro Asn Gly Ser Val Pro Gly Asp Arg Ile Thr Phe Asp Ala Phe Pro  
115 120 125  
Gly Glu Pro Asp Lys Glu Leu Asn Pro Lys Lys Lys Ile Trp Glu Gln  
130 135 140  
Ile Gln Pro Asp Leu His Thr Asn Asp Glu Cys Val Ala Thr Tyr Lys  
145 150 155 160  
Gly Val Pro Phe Glu Val Lys Gly Lys Gly Val Cys Arg Ala Gln Thr  
165 170 175  
Met Ser Asn Ser Gly Ile Lys  
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*JZ*  
*Cmt*

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